

SEQUENCE LISTING

<110> Conklin, Darrell C.
Haldeman, Betty A.

<120> MAMMALIAN CYTOKINE-LIKE POLYPEPTIDE-10

<130> 97-72

<150> 09/199,586

<151> 1998-11-25

<150> 60/066,597

<151> 1997-11-26

<160> 43

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 926

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (45)...(572)

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agt ctt gcc ttc agc ctt ctc tct gct gcg ttt tat ctc cta tgg act 104
Ser Leu Ala Phe Ser Leu Leu Ser Ala Ala Phe Tyr Leu Leu Trp Thr
5 10 15 20

cct tcc act gga ctg aag aca ctc aat ttg gga agc tgt gtg atc gcc 152
Pro Ser Thr Gly Leu Lys Thr Leu Asn Leu Gly Ser Cys Val Ile Ala
25 30 35

aca aac ctt cag gaa ata cga aat gga ttt tct gac ata cgg ggc agt 200

Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp Ile Arg Gly Ser	
40 45 50	
gtg caa gcc aaa gat gga aac att gac atc aga atc tta agg agg act	248
Val Gln Ala Lys Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg Arg Thr	
55 60 65	
gag tct ttg caa gac aca aag cct gcg aat cga tgc tgc ctc ctg cgc	296
Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu Leu Arg	
70 75 80	
cat ttg cta aga ctc tat ctg gac agg gta ttt aaa aac tac cag acc	344
His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr	
85 90 95 100	
cct gac cat tat act ctc cgg aag atc agc agc ctc gcc aat tcc ttt	392
Pro Asp His Tyr Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe	
105 110 115	
ctt acc atc aag aag gac ctc cgg ctc tgt cat gcc cac atg aca tgc	440
Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys His Ala His Met Thr Cys	
120 125 130	
cat tgt ggg gag gaa gca atg aag aaa tac agc cag att ctg agt cac	488
His Cys Gly Glu Glu Ala Met Lys Lys Tyr Ser Gln Ile Leu Ser His	
135 140 145	
ttt gaa aag ctg gaa cct cag gca gca gtt gtg aag gct ttg ggg gaa	536
Phe Glu Lys Leu Glu Pro Gln Ala Ala Val Val Lys Ala Leu Gly Glu	
150 155 160	
cta gac att ctt ctg caa tgg atg gag gag aca gaa taggaggaaa	582
Leu Asp Ile Leu Leu Gln Trp Met Glu Glu Thr Glu	
165 170 175	
gtgatgctgc tgctaagaat attcgaggtc aagagctcca gtcttcaata cctgcagagg	642
aggcatgacc ccaaaccacc atctctttac tgtactagtc ttgtgctggt cacagtgtat	702
cttattttatg cattacttgc ttccttgcac gattgtcttt atgcatcccc aatcttaatt	762
gagaccatac ttgtataaga tttttgtaat atctttctgc tattggatat atttattagt	822
taatataattt atttattttt tgctattaat gtattttaatt ttttacttgg gcatgaaact	882
ttaaaaaaaaaa ttcacaagat tatatttata acctgactag agca	926

<211> 176

<212> PRT

<213> Homo sapiens

<400> 2

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Met Lys Ala Ser Ser Leu Ala Phe Ser Leu Leu Ser Ala Ala Phe Tyr
 1           5           10           15
Leu Leu Trp Thr Pro Ser Thr Gly Leu Lys Thr Leu Asn Leu Gly Ser
 20           25           30
Cys Val Ile Ala Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp
 35           40           45
Ile Arg Gly Ser Val Gln Ala Lys Asp Gly Asn Ile Asp Ile Arg Ile
 50           55           60
Leu Arg Arg Thr Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys
 65           70           75           80
Cys Leu Leu Arg His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys
 85           90           95
Asn Tyr Gln Thr Pro Asp His Tyr Thr Leu Arg Lys Ile Ser Ser Leu
100           105           110
Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys His Ala
115           120           125
His Met Thr Cys His Cys Gly Glu Glu Ala Met Lys Lys Tyr Ser Gln
130           135           140
Ile Leu Ser His Phe Glu Lys Leu Glu Pro Gln Ala Ala Val Val Lys
145           150           155           160
Ala Leu Gly Glu Leu Asp Ile Leu Leu Gln Trp Met Glu Glu Thr Glu
165           170           175

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<210> 3

<211> 793

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (45)...(497)

<400> 3

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                               Met Lys Ala Ser
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agt ctt gcc ttc agc ctt ctc tct gct gcg ttt tat ctc cta tgg act      104

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gtattttaatt ttttac

793

<210> 4

<211> 151

<212> PRT

<213> Homo sapiens

<400> 4

Met	Lys	Ala	Ser	Ser	Leu	Ala	Phe	Ser	Leu	Leu	Ser	Ala	Ala	Phe	Tyr
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Leu	Leu	Trp	Thr	Pro	Ser	Thr	Gly	Leu	Lys	Thr	Leu	Asn	Leu	Gly	Ser
		20					25					30			
Cys	Val	Ile	Ala	Thr	Asn	Leu	Gln	Glu	Ile	Arg	Asn	Gly	Phe	Ser	Asp
		35				40						45			
Ile	Arg	Gly	Ser	Val	Gln	Ala	Lys	Asp	Gly	Asn	Ile	Asp	Ile	Arg	Ile
	50				55					60					
Leu	Arg	Arg	Thr	Glu	Ser	Leu	Gln	Asp	Thr	Lys	Pro	Ala	Asn	Arg	Cys
65				70				75						80	
Cys	Leu	Leu	Arg	His	Leu	Leu	Arg	Leu	Tyr	Leu	Asp	Arg	Val	Phe	Lys
			85					90					95		
Asn	Tyr	Gln	Thr	Pro	Asp	His	Tyr	Thr	Leu	Arg	Lys	Ile	Ser	Ser	Leu
		100					105					110			
Ala	Asn	Ser	Phe	Leu	Thr	Ile	Lys	Lys	Asp	Leu	Arg	Leu	Cys	Leu	Glu
	115					120				125					
Pro	Gln	Ala	Ala	Val	Val	Lys	Ala	Leu	Gly	Glu	Leu	Asp	Ile	Leu	Leu
	130					135				140					
Gln	Trp	Met	Glu	Glu	Thr	Glu									
145					150										

<210> 5

<211> 253

<212> DNA

<213> Homo sapiens

<400> 5

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agacactcaa	tttggaagc	tgtgtgatcg	ccacaaacct	tcaggaaata	cgaaatggat	180
tttctgagat	acggggcagt	gtgcaagcca	aagatggaaa	cattgacatc	agaatcttaa	240
ggaggactga	gtc					253

<210> 6

<211> 24

<212> DNA

<213> Homo sapiens

<400> 6

attcctagct cctgtggtct ccag

24

<210> 7

<211> 25

<212> DNA

<213> Homo sapiens

<400> 7

ctctgctgcg ttttatctcc tatgg

25

<210> 8

<211> 22

<212> DNA

<213> Homo sapiens

<400> 8

tcccaaattg agtgtcttca gt

22

<210> 9

<211> 45

<212> DNA

<213> Homo sapiens

<400> 9

cacagcttcc caaattgagt gtcttcagtc cagtggaagg agtcc

45

<210> 10

<211> 747

<212> DNA

<213> Homo sapiens

<400> 10

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ttgctaagac tctatctgga cagggtattht aaaaactacc agaccctga ccattatact	180
ctccggaaga tcagcagcct cgccaattcc tttcttacca tcaagaagga cctccggctc	240
tgtcatgccc acatgacatg ccattgtggg gaggaagcaa tgaagaaata cagccagatt	300
ctgagtcact ttgaaaagct ggaacctcag gcagcagttg tgaaggcttt gggggaacta	360
gacattcttc tgcaatggat ggaggagaca gaataggagg aaagtgatgc tgctgctaag	420

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aatattcgag gtcaagagct ccagtcttca atacctgcag aggaggcatg accccaaacc 480
accatctctt tactgtacta gtcttgtgct ggtcacagtg tatcttattt atgcattact 540
tgcttccttg catgattgtc tttatgcata cccaatctta attgagacca tacttgtata 600
agatTTTTgt aatatctttc tgctattgga tatatttatt agttaatata tttatttatt 660
ttttgctatt aatgtattta attttttact tgggcatgaa actttaaaaa aaattcacia 720
gattatattt ataacctgac tagagca 747

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<210> 11
<211> 614
<212> DNA
<213> Homo sapiens

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<400> 11
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aggaggactg agtcttttga agacacaaag cctgcgaatc gatgctgcct cctgcgccat 120
ttgctaagac tctatctgga cagggtattt aaaaactacc agaccctga ccattatact 180
ctccggaaga tcagcagcct cgccaattcc tttcttacca tcaagaagga cctccggctc 240
tgtctggaac ctcaggcagc agttgtgaag gctttggggg aactagacat tcttctgcaa 300
tggatggagg agacagaata ggaggaaagt gatgctgctg ctaagaatat tcgagggtcaa 360
gagctccagt cttcaatacc tgcagaggag gcatgacccc aaaccacat ctctttactg 420
tactagtctt gtgctgggtca cagtgtatct tatttatgca ttacttgctt ccttgcata 480
ttgtctttat gcatcccaa tcttaattga gaccatactt gtataagatt tttgtaatat 540
ctttctgcta ttggatatat ttattagtta atatatttat ttattttttg ctattaatgt 600
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<210> 12
<211> 152
<212> PRT
<213> Homo sapiens

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<400> 12
Leu Lys Thr Leu Asn Leu Gly Ser Cys Val Ile Ala Thr Asn Leu Gln
 1          5          10          15
Glu Ile Arg Asn Gly Phe Ser Asp Ile Arg Gly Ser Val Gln Ala Lys
 20          25          30
Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg Arg Thr Glu Ser Leu Gln
 35          40          45
Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu Leu Arg His Leu Leu Arg
 50          55          60
Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr Pro Asp His Tyr
 65          70          75          80
Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Thr Ile Lys
          85          90          95

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Lys Asp Leu Arg Leu Cys His Ala His Met Thr Cys His Cys Gly Glu
 100 105 110
 Glu Ala Met Lys Lys Tyr Ser Gln Ile Leu Ser His Phe Glu Lys Leu
 115 120 125
 Glu Pro Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu Asp Ile Leu
 130 135 140
 Leu Gln Trp Met Glu Glu Thr Glu
 145 150

<210> 13
 <211> 127
 <212> PRT
 <213> Homo sapiens

<400> 13

Leu Lys Thr Leu Asn Leu Gly Ser Cys Val Ile Ala Thr Asn Leu Gln
 1 5 10 15
 Glu Ile Arg Asn Gly Phe Ser Asp Ile Arg Gly Ser Val Gln Ala Lys
 20 25 30
 Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg Arg Thr Glu Ser Leu Gln
 35 40 45
 Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu Leu Arg His Leu Leu Arg
 50 55 60
 Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr Pro Asp His Tyr
 65 70 75 80
 Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Thr Ile Lys
 85 90 95
 Lys Asp Leu Arg Leu Cys Leu Glu Pro Gln Ala Ala Val Val Lys Ala
 100 105 110
 Leu Gly Glu Leu Asp Ile Leu Leu Gln Trp Met Glu Glu Thr Glu
 115 120 125

<210> 14
 <211> 15
 <212> PRT
 <213> Homo sapiens

<400> 14

Ile Ala Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp Ile
 1 5 10 15

<210> 15
 <211> 15

<213> Homo sapiens

<400> 15

Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr Pro Asp His Tyr Thr
1 5 10 15

<210> 16

<211> 15

<212> PRT

<213> Homo sapiens

<400> 16

Leu Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys
1 5 10 15

<210> 17

<211> 15

<212> PRT

<213> Homo sapiens

<400> 17

Val Val Lys Ala Leu Gly Glu Leu Asp Ile Leu Leu Gln Trp Met
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<210> 18

<211> 824

<212> DNA

<213> Mus musculus

 $\langle 220 \rangle$

<221> CDS

<222> (71)...(598)

<400> 18

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      Met Lys Gly Phe Gly Leu Ala Phe Gly Leu Phe Ser Ala
          1              5              10

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gtg ggt ttt ctt ctc tgg act cct tta act ggg ctc aag acc ctc cat 157
Val Gly Phe Leu Leu Trp Thr Pro Leu Thr Gly Leu Lys Thr Leu His
15 20 25

ttg gga agc tgt gtg att act gca aac cta cag gca ata caa aag gaa Leu Gly Ser Cys Val Ile Thr Ala Asn Leu Gln Ala Ile Gln Lys Glu 30 35 40 45	205
ttt tct gag att cgg gat agt gtg caa gct gaa gat aca aat att gac Phe Ser Glu Ile Arg Asp Ser Val Gln Ala Glu Asp Thr Asn Ile Asp 50 55 60	253
atc aga att tta agg acg act gag tct ttg aaa gac ata aag tct ttg Ile Arg Ile Leu Arg Thr Thr Glu Ser Leu Lys Asp Ile Lys Ser Leu 65 70 75	301
gat agg tgc tgc ttc ctt cgt cat cta gtg aga ttc tat ctg gac agg Asp Arg Cys Cys Phe Leu Arg His Leu Val Arg Phe Tyr Leu Asp Arg 80 85 90	349
gta ttc aaa gtc tac cag acc cct gac cac cat acc ctg aga aag atc Val Phe Lys Val Tyr Gln Thr Pro Asp His His Thr Leu Arg Lys Ile 95 100 105	397
agc agc ctc gcc aac tcc ttt ctt atc atc aag aag gac ctc tca gtc Ser Ser Leu Ala Asn Ser Phe Leu Ile Ile Lys Lys Asp Leu Ser Val 110 115 120 125	445
tgt cat tct cac atg gca tgt cat tgt ggg gaa gaa gca atg gag aaa Cys His Ser His Met Ala Cys His Cys Gly Glu Glu Ala Met Glu Lys 130 135 140	493
tac aac caa att ctg agt cac ttc ata gag ttg gaa ctt cag gca gcg Tyr Asn Gln Ile Leu Ser His Phe Ile Glu Leu Glu Leu Gln Ala Ala 145 150 155	541
gtg gta aag gct ttg gga gaa cta ggc att ctt ctg aga tgg atg gag Val Val Lys Ala Leu Gly Glu Leu Gly Ile Leu Leu Arg Trp Met Glu 160 165 170	589
gag atg cta tagatgaaag tggagaggct gctgagaaca ctcctgtcca Glu Met Leu 175	638
agaatctcag acctcagcac catgaagaca tggccccagg tgctggcatt tctactcaag	698

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agttccagtc ctcagcacca cgaagatggc ctcaaaccac caccctttg tgatataact 758
tagtgctagc tatgtgtata ttatttctac attattggct cccttatgtg aatgccttca 818
tgtgtc 824

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<210> 19
<211> 176
<212> PRT
<213> Mus musculus

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<400> 19
Met Lys Gly Phe Gly Leu Ala Phe Gly Leu Phe Ser Ala Val Gly Phe
 1          5          10          15
Leu Leu Trp Thr Pro Leu Thr Gly Leu Lys Thr Leu His Leu Gly Ser
 20          25          30
Cys Val Ile Thr Ala Asn Leu Gln Ala Ile Gln Lys Glu Phe Ser Glu
 35          40          45
Ile Arg Asp Ser Val Gln Ala Glu Asp Thr Asn Ile Asp Ile Arg Ile
 50          55          60
Leu Arg Thr Thr Glu Ser Leu Lys Asp Ile Lys Ser Leu Asp Arg Cys
65 70          75          80
Cys Phe Leu Arg His Leu Val Arg Phe Tyr Leu Asp Arg Val Phe Lys
 85          90          95
Val Tyr Gln Thr Pro Asp His His Thr Leu Arg Lys Ile Ser Ser Leu
100          105          110
Ala Asn Ser Phe Leu Ile Ile Lys Lys Asp Leu Ser Val Cys His Ser
115          120          125
His Met Ala Cys His Cys Gly Glu Glu Ala Met Glu Lys Tyr Asn Gln
130          135          140
Ile Leu Ser His Phe Ile Glu Leu Glu Leu Gln Ala Ala Val Val Lys
145          150          155          160
Ala Leu Gly Glu Leu Gly Ile Leu Leu Arg Trp Met Glu Glu Met Leu
165          170          175

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<210> 20
<211> 152
<212> PRT
<213> Mus musculus

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<400> 20
Leu Lys Thr Leu His Leu Gly Ser Cys Val Ile Thr Ala Asn Leu Gln
 1          5          10          15
Ala Ile Gln Lys Glu Phe Ser Glu Ile Arg Asp Ser Val Gln Ala Glu
 20          25          30

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1. 1. 1. 1.

<210> 24
 <211> 15
 <212> PRT
 <213> Mus musculus

<400> 24
 Val Val Lys Ala Leu Gly Glu Leu Gly Ile Leu Leu Arg Trp Met
 1 5 10 15

<210> 25
 <211> 144
 <212> PRT
 <213> Mus musculus

<400> 25
 Cys Val Ile Thr Ala Asn Leu Gln Ala Ile Gln Lys Glu Phe Ser Glu
 1 5 10 15
 Ile Arg Asp Ser Val Gln Ala Glu Asp Thr Asn Ile Asp Ile Arg Ile
 20 25 30
 Leu Arg Thr Thr Glu Ser Leu Lys Asp Ile Lys Ser Leu Asp Arg Cys
 35 40 45
 Cys Phe Leu Arg His Leu Val Arg Phe Tyr Leu Asp Arg Val Phe Lys
 50 55 60
 Val Tyr Gln Thr Pro Asp His His Thr Leu Arg Lys Ile Ser Ser Leu
 65 70 75 80
 Ala Asn Ser Phe Leu Ile Ile Lys Lys Asp Leu Ser Val Cys His Ser
 85 90 95
 His Met Ala Cys His Cys Gly Glu Glu Ala Met Glu Lys Tyr Asn Gln
 100 105 110
 Ile Leu Ser His Phe Ile Glu Leu Glu Leu Gln Ala Ala Val Val Lys
 115 120 125
 Ala Leu Gly Glu Leu Gly Ile Leu Leu Arg Trp Met Glu Glu Met Leu
 130 135 140

<210> 26
 <211> 144
 <212> PRT
 <213> Homo sapiens

<400> 26
 Cys Val Ile Ala Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp
 1 5 10 15

Ile Arg Gly Ser Val Gln Ala Lys Asp Gly Asn Ile Asp Ile Arg Ile
 20 25 30
 Leu Arg Arg Thr Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys
 35 40 45
 Cys Leu Leu Arg His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys
 50 55 60
 Asn Tyr Gln Thr Pro Asp His Tyr Thr Leu Arg Lys Ile Ser Ser Leu
 65 70 75 80
 Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys His Ala
 85 90 95
 His Met Thr Cys His Cys Gly Glu Glu Ala Met Lys Lys Tyr Ser Gln
 100 105 110
 Ile Leu Ser His Phe Glu Lys Leu Glu Pro Gln Ala Ala Val Val Lys
 115 120 125
 Ala Leu Gly Glu Leu Asp Ile Leu Leu Gln Trp Met Glu Glu Thr Glu
 130 135 140

<210> 27

<211> 38

<212> PRT

<213> Homo sapiens

<400> 27

Cys Gly Glu Glu Ala Met Lys Lys Tyr Ser Gln Ile Leu Ser His Phe
 1 5 10 15

Glu Lys Leu Glu Pro Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu
 20 25 30
 Asp Ile Leu Leu Gln Trp
 35

<210> 28

<211> 71

<212> PRT

<213> Homo sapiens

<400> 28

Ile Ala Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp Ile Arg
 1 5 10 15
 Gly Ser Val Gln Ala Lys Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg
 20 25 30
 Arg Thr Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu
 35 40 45

Leu Arg His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr
 50 55 60
 Gln Thr Pro Asp His Tyr Thr
 65 70

<210> 29
 <211> 92
 <212> PRT
 <213> Homo sapiens

<400> 29
 Ile Ala Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp Ile Arg
 1 5 10 15
 Gly Ser Val Gln Ala Lys Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg
 20 25 30
 Arg Thr Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu
 35 40 45
 Leu Arg His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr
 50 55 60
 Gln Thr Pro Asp His Tyr Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn
 65 70 75 80
 Ser Phe Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys
 85 90

<210> 30
 <211> 82
 <212> PRT
 <213> Homo sapiens

<400> 30
 Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr Pro Asp His Tyr Thr Leu
 1 5 10 15
 Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp
 20 25 30
 Leu Arg Leu Cys His Ala His Met Thr Cys His Cys Gly Glu Glu Ala
 35 40 45
 Met Lys Lys Tyr Ser Gln Ile Leu Ser His Phe Glu Lys Leu Glu Pro
 50 55 60
 Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu Asp Ile Leu Leu Gln
 65 70 75 80
 Trp Met

<210> 31

<211> 36
 <212> PRT
 <213> Homo sapiens

<400> 31
 Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr Pro Asp His Tyr Thr Leu
 1 5 10 15
 Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp
 20 25 30
 Leu Arg Leu Cys
 35

<210> 32
 <211> 61
 <212> PRT
 <213> Homo sapiens

<400> 32
 Leu Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys His
 1 5 10 15
 Ala His Met Thr Cys His Cys Gly Glu Glu Ala Met Lys Lys Tyr Ser
 20 25 30
 Gln Ile Leu Ser His Phe Glu Lys Leu Glu Pro Gln Ala Ala Val Val
 35 40 45
 Lys Ala Leu Gly Glu Leu Asp Ile Leu Leu Gln Trp Met
 50 55 60

<210> 33
 <211> 756
 <212> DNA
 <213> Mus musculus

<220>
 <221> CDS
 <222> (71)...(532)

<400> 33
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 taggtgtaag atg aaa ggc ttt ggt ctt gcc ttt gga ctg ttc tcc gct 109
 Met Lys Gly Phe Gly Leu Ala Phe Gly Leu Phe Ser Ala
 1 5 10
 gtg ggt ttt ctt ctc tgg act cct tta act ggg ctc aag acc ctc cat 157

Val Gly Phe Leu Leu Trp Thr Pro Leu Thr Gly Leu Lys Thr Leu His	
15 20 25	
ttg gga agc tgt gtg att act gca aac cta cag gca ata caa aag gaa	205
Leu Gly Ser Cys Val Ile Thr Ala Asn Leu Gln Ala Ile Gln Lys Glu	
30 35 40 45	
ttt tct gag att cgg gat agt gtg tct ttg gat agg tgc tgc ttc ctt	253
Phe Ser Glu Ile Arg Asp Ser Val Ser Leu Asp Arg Cys Cys Phe Leu	
50 55 60	
cgt cat cta gtg aga ttc tat ctg gac agg gta ttc aaa gtc tac cag	301
Arg His Leu Val Arg Phe Tyr Leu Asp Arg Val Phe Lys Val Tyr Gln	
65 70 75	
acc cct gac cac cat acc ctg aga aag atc agc agc ctc gcc aac tcc	349
Thr Pro Asp His His Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser	
80 85 90	
ttt ctt atc atc aag aag gac ctc tca gtc tgt cat tct cac atg gca	397
Phe Leu Ile Ile Lys Lys Asp Leu Ser Val Cys His Ser His Met Ala	
95 100 105	
tgt cat tgt ggg gaa gaa gca atg gag aaa tac aac caa att ctg agt	445
Cys His Cys Gly Glu Glu Ala Met Glu Lys Tyr Asn Gln Ile Leu Ser	
110 115 120 125	
cac ttc ata gag ttg gaa ctt cag gca gcg gtg gta aag gct ttg gga	493
His Phe Ile Glu Leu Glu Leu Gln Ala Ala Val Val Lys Ala Leu Gly	
130 135 140	
gaa cta ggc att ctt ctg aga tgg atg gag gag atg cta tagatgaaag	542
Glu Leu Gly Ile Leu Leu Arg Trp Met Glu Glu Met Leu	
145 150	
tggaataggct gctgagaaca ctctgtcca agaatctcag acctcagcac catgaagaca	602
tggtcccagg tgctggcatt tctactcaag agttccagtc ctcagcacca cgaagatggc	662
ctcaaaccac caccctttg tgatataact tagtgctagc tatgtgtata ttatttctac	722
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<213> Mus musculus

<400> 34

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Met Lys Gly Phe Gly Leu Ala Phe Gly Leu Phe Ser Ala Val Gly Phe
 1           5           10           15
Leu Leu Trp Thr Pro Leu Thr Gly Leu Lys Thr Leu His Leu Gly Ser
 20           25           30
Cys Val Ile Thr Ala Asn Leu Gln Ala Ile Gln Lys Glu Phe Ser Glu
 35           40           45
Ile Arg Asp Ser Val Ser Leu Asp Arg Cys Cys Phe Leu Arg His Leu
 50           55           60
Val Arg Phe Tyr Leu Asp Arg Val Phe Lys Val Tyr Gln Thr Pro Asp
 65           70           75           80
His His Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Ile
 85           90           95
Ile Lys Lys Asp Leu Ser Val Cys His Ser His Met Ala Cys His Cys
 100          105          110
Gly Glu Glu Ala Met Glu Lys Tyr Asn Gln Ile Leu Ser His Phe Ile
 115          120          125
Glu Leu Glu Leu Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu Gly
 130          135          140
Ile Leu Leu Arg Trp Met Glu Glu Met Leu
145          150

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<210> 35

<211> 130

<212> PRT

<213> Mus musculus

<400> 35

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Leu Lys Thr Leu His Leu Gly Ser Cys Val Ile Thr Ala Asn Leu Gln
 1           5           10           15

Ala Ile Gln Lys Glu Phe Ser Glu Ile Arg Asp Ser Val Ser Leu Asp
 20           25           30
Arg Cys Cys Phe Leu Arg His Leu Val Arg Phe Tyr Leu Asp Arg Val
 35           40           45
Phe Lys Val Tyr Gln Thr Pro Asp His His Thr Leu Arg Lys Ile Ser
 50           55           60
Ser Leu Ala Asn Ser Phe Leu Ile Ile Lys Lys Asp Leu Ser Val Cys
 65           70           75           80
His Ser His Met Ala Cys His Cys Gly Glu Glu Ala Met Glu Lys Tyr

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				85					90					95	
Asn	Gln	Ile	Leu	Ser	His	Phe	Ile	Glu	Leu	Glu	Leu	Gln	Ala	Ala	Val
			100					105					110		
Val	Lys	Ala	Leu	Gly	Glu	Leu	Gly	Ile	Leu	Leu	Arg	Trp	Met	Glu	Glu
		115					120					125			
Met	Leu														
	130														

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26

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23

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25